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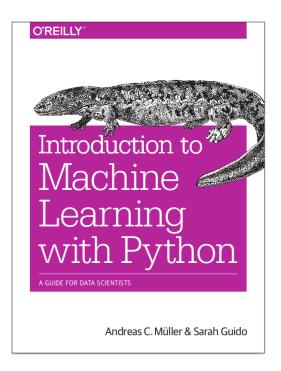






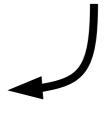


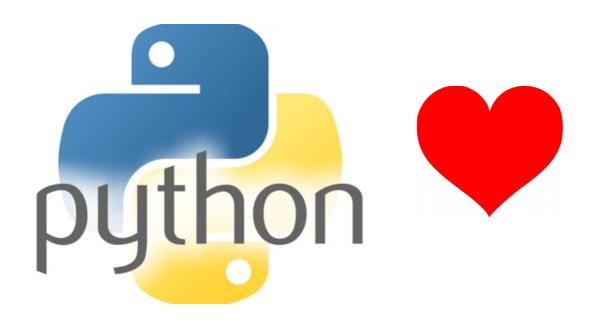














Classification Regression Clustering Semi-Supervised Learning Feature Selection Feature Extraction Manifold Learning **Dimensionality Reduction** Kernel Approximation Hyperparameter Optimization **Evaluation Metrics** Out-of-core learning

















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sklearn-wheels



Tom Dupré la Tour





Virgile Fritsch VirgileFritsch



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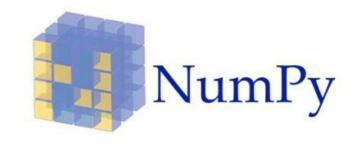


#### Mission

Commoditize and Democratize Machine Learning

#### **Basic API**

# Representing Data



1.1	2.2	3.4	5.6	1.0	$\setminus$
6.7	0.5	0.4	2.6	1.6	
2.4	9.3	7.3	6.4	2.8	
1.5	0.0	4.3	8.3	3.4	
0.5	3.5	8.1	3.6	4.6	
5.1	9.7	3.5	7.9	5.1	
3.7	7.8	2.6	3.2	6.3	
	6.7 2.4 1.5 0.5 5.1	6.7 0.5 2.4 9.3 1.5 0.0 0.5 3.5 5.1 9.7	6.70.50.42.49.37.31.50.04.30.53.58.15.19.73.5	6.7       0.5       0.4       2.6         2.4       9.3       7.3       6.4         1.5       0.0       4.3       8.3         0.5       3.5       8.1       3.6         5.1       9.7       3.5       7.9	2.4       9.3       7.3       6.4       2.8         1.5       0.0       4.3       8.3       3.4         0.5       3.5       8.1       3.6       4.6         5.1       9.7       3.5       7.9       5.1

one feature

outputs / labels

clf = RandomForestClassifier() clf.fit(X\_train, y\_train) Training Data Model Training Labels y\_pred = clf.predict(X\_test) Prediction **Test Data** clf.score(X\_test, y\_test) **Test Labels Evaluation** 

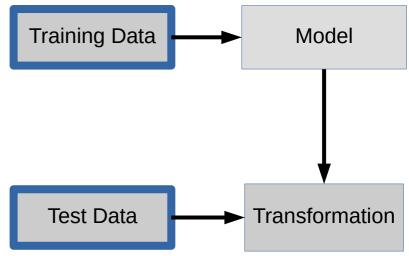
### **Unsupervised Transformations**

```
pca = PCA()

pca.fit(X_train)

Training Data

X_new = pca.transform(X_test)
```



### Core API Summary

estimator.fit(X, [y])

estimator.predict

estimator.transform

Classification

Preprocessing

Regression

Dimensionality reduction

Clustering

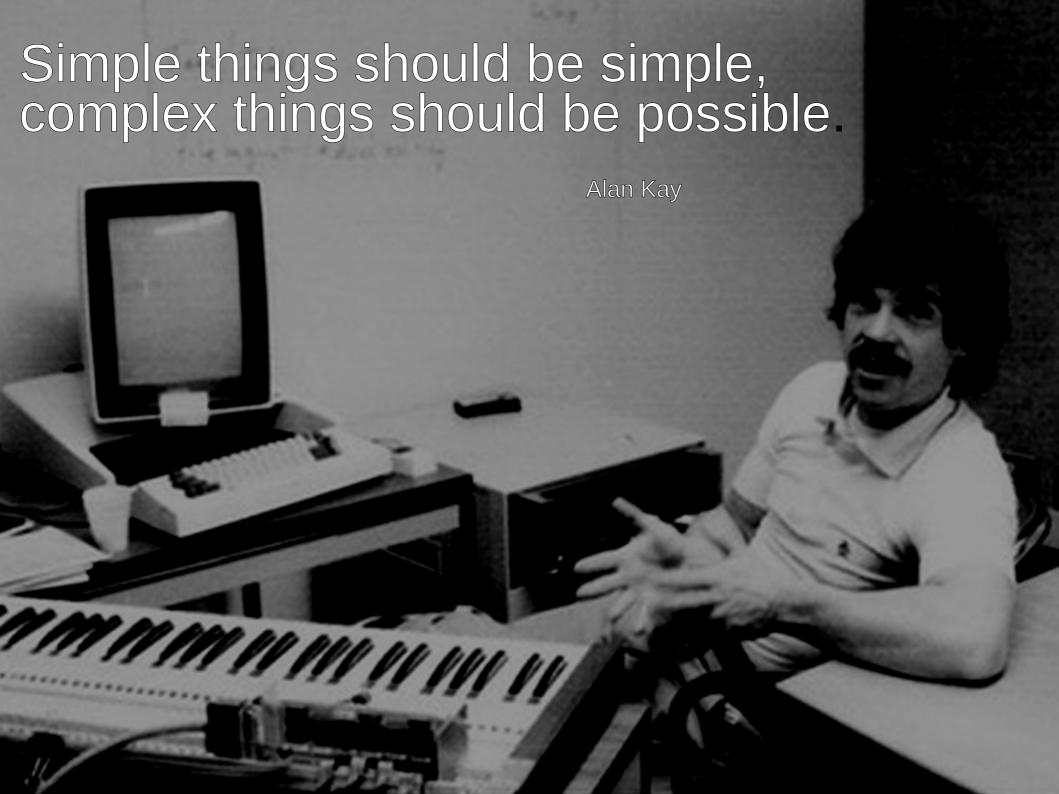
Feature selection

Feature extraction

#### **Guiding Ideas**

Goals:

Maintainability Ease of use



# Three way documentation

#### 1.9. Ensemble methods

The goal of **ensemble methods** is to combine the predictions of several base estimators built with a given learning algorithm in order to improve generalizability / robustness over a single estimator.

Two families of ensemble methods are usually distinguished:

 In averaging methods, the driving principle is to build several estimators independently and then to average their predictions. On average, the combined estimator is usually better than any of the single base estimator because its variance is reduced.

**Examples:** Bagging methods, Forests of randomized trees, ...

 By contrast, in boosting methods, base estimators are built sequentially and one tries to reduce the bias of the combined estimator. The motivation is to combine several weak models to produce a powerful ensemble.

Examples: AdaBoost, Gradient Tree Boosting, ...

#### sklearn.ensemble.RandomForestClassifier

class sklearn.ensemble. RandomForestClassifier (n\_estimators=10, criterion='gini', max\_depth=None, min\_samples\_split=2, min\_samples\_leaf=1, min\_weight\_fraction\_leaf=0.0, max\_features='auto', max\_leaf\_nodes=None, bootstrap=True, oob\_score=False, n\_jobs=1, random\_state=None, verbose=0, warm\_start=False) [source]

A random forest classifier.

A random forest is a meta estimator that fits a number of decision tree classifiers on various sub-samples of the dataset and use averaging to improve the predictive accuracy and control over-fitting.

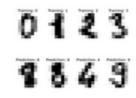
Parameters: n estimators: integer, optional (default=10)

The number of trees in the forest.

criterion: string, optional (default="gini")

The function to measure the quality of a split. Supported criteria are "gini" for the Gini impurity and "entropy" for the information gain. Note: this parameter is tree-specific.

#### **Examples**



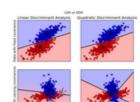
Recognizing hand-written digits



Plot classification probability



Classifier comparison



Linear and Quadratic
Discriminant Analysis

with confidence ellipsoid

#### Ease of Installation

# Simplicity

```
lr = LogisticRegression()
lr.fit(X_train, y_train)
lr.score(X_test, y_test)
```

### Consistency

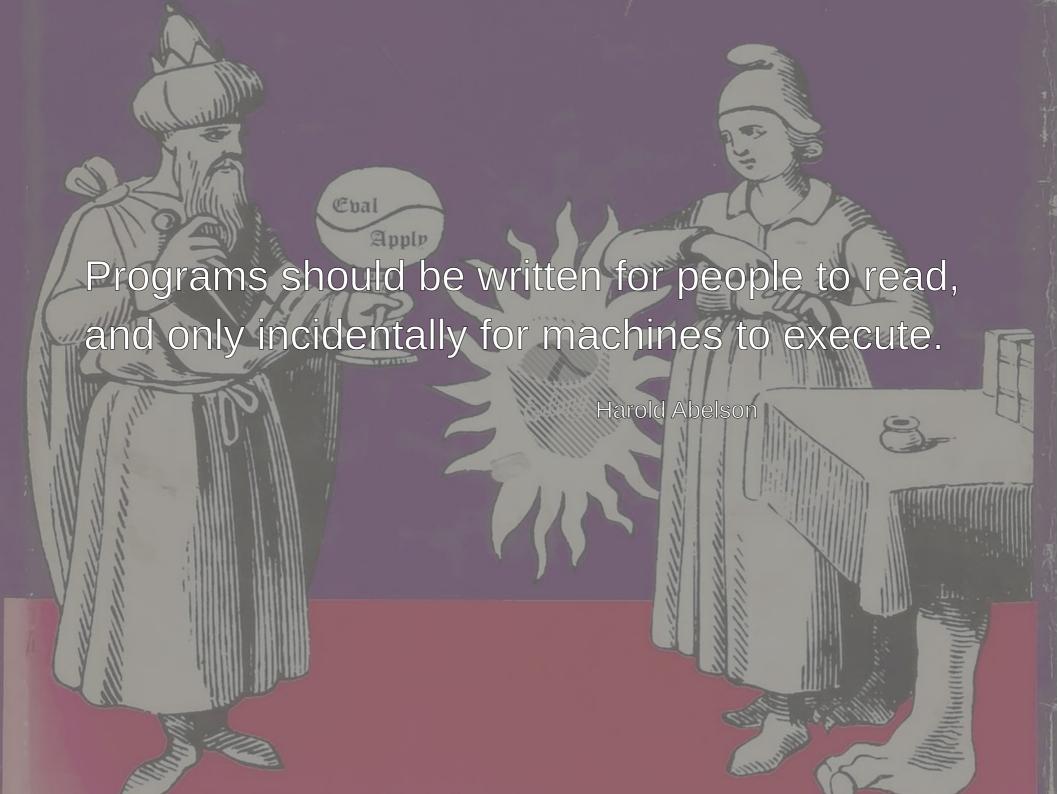
```
grid = GridSearchCV(svm,param_grid)
grid.fit(X_train, y_train)
grid.score(X_test, y_test)
```

# Composition

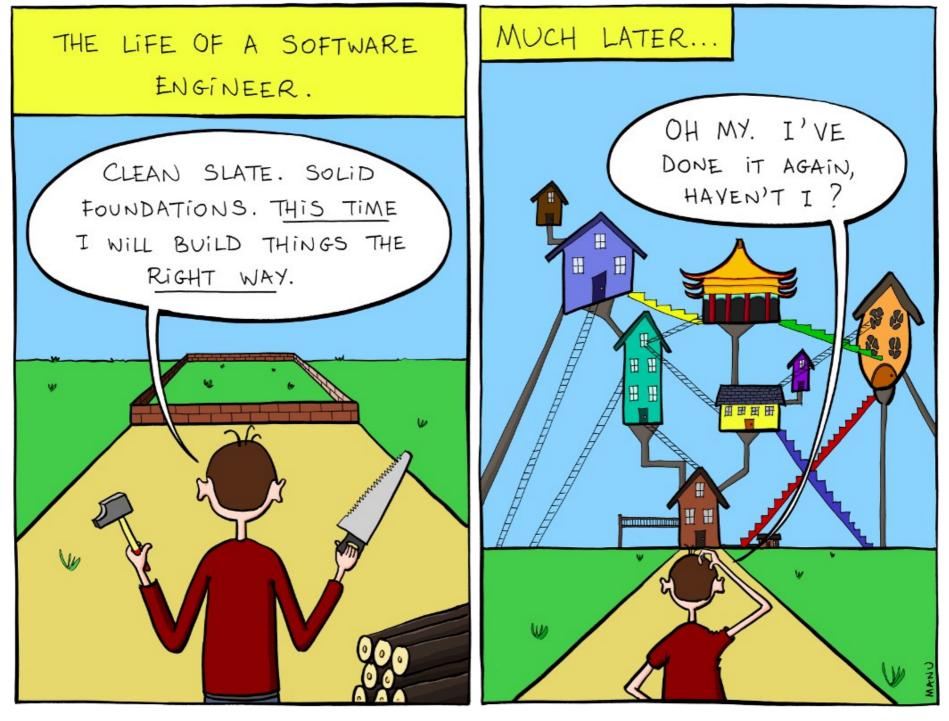
#### Default Parameters

### Flat Class Hierarchy, Few Types

- Numpy arrays / sparse matrices
- Estimators
- [Cross-validation objects]
- [Scorers]

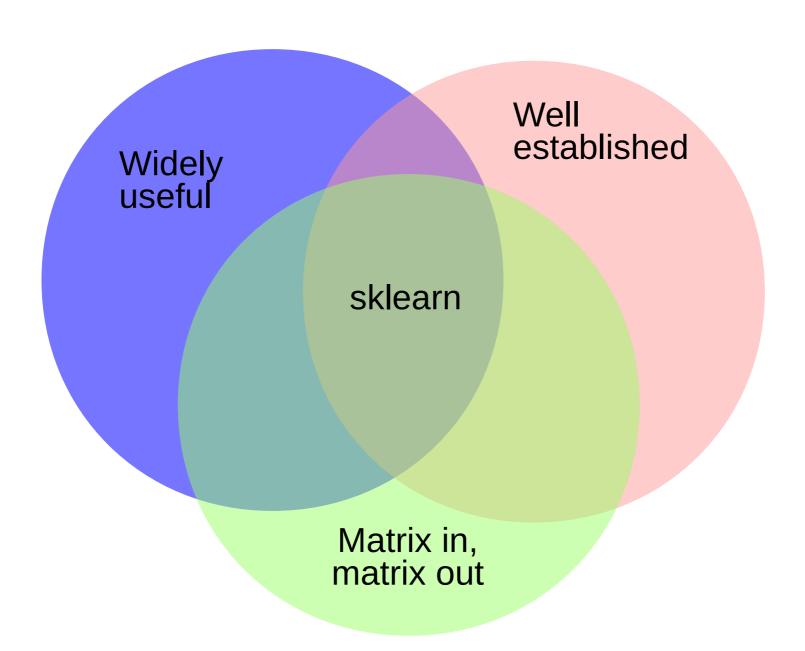






Feature Creep

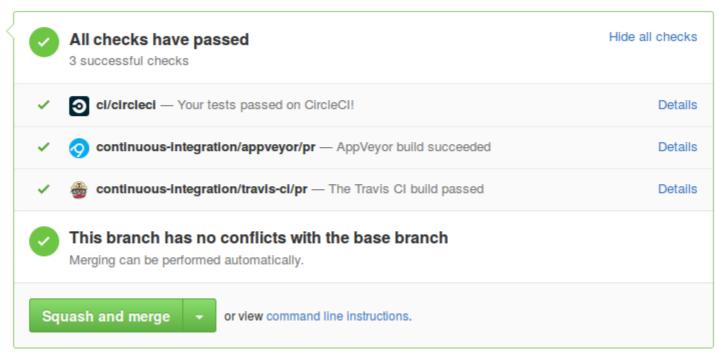
# Scoping



# Testing & Continuous Integration

Add more commits by pushing to the more\_repr branch on amueller/scikit-learn.







#### API implementation & contracts

# Method Chaining

Fit must return self:

#### fit resets

```
from sklearn.tree import DecisionTreeClassifier
tree = DecisionTreeClassifier(max_depth=10)
tree.fit(iris.data, iris.target)
score_iris = tree.score(iris.data, iris.target)
tree.fit(digits.data, digits.target)
score_digits = tree.score(digits.data, digits.target)
```

# (only) fit mutates self

- \_\_init\_\_ only remembers construction parameters
- Transform / score etc don't change object
- Fit returns self but mutates object!

#### Estimated attributes

```
Attributes
 components : array, shape (n components, n features)
     Principal axes in feature space, representing the directions of
     maximum variance in the data. The components are sorted by
     ``explained variance ``.
 explained variance : array, shape (n components,)
     The amount of variance explained by each of the selected components.
     Equal to n components largest eigenvalues
     of the covariance matrix of X.
     .. versionadded:: 0.18
 explained variance ratio : array, shape (n components,)
     Percentage of variance explained by each of the selected components.
     If ``n components`` is not set then all components are stored and the
     sum of the ratios is equal to 1.0.
 singular values : array, shape (n components,)
     The singular values corresponding to each of the selected components.
     The singular values are equal to the 2-norms of the ``n components``
     variables in the lower-dimensional space.
Leaves two kinds of attributes:
arguments to init; things estimated during fit
```

# fit\_transform / fit\_predict

In general: computational shortcut:

```
pca = PCA()
pca.fit(X)
X_pca = pca.transform(X)
X_pca2 = pca.fit_transform(X)
```

Clustering / Manifold learning: Not inductive.

```
tsne = TSNE()
X_tsne = tsne.fit_transform(X)
```

```
dbscan = DBSCAN()
cluster_labels = dbscan.fit_predict(X)
```

### check\_estimator

```
class TemplateClassifier(BaseEstimator, ClassifierMixin):
   def init (self, demo param='demo'):
        self.demo param = demo param
   def fit(self, X, y):
       # Check that X and y have correct shape
       X, y = \text{check } X y(X, y)
       # Store the classes seen during fit
       self.classes = unique labels(y)
        self.X = X
        self.v = v
       # Return the classifier
        return self
   def predict(self, X):
        closest = np.argmin(euclidean distances(X, self.X), axis=1)
        return self.v [closest]
```

#### check\_estimator(TemplateClassifier)

AssertionError: Error message does not include the expected string: 'fit'. Observed error message: "'TemplateClassifier' object has no attribute 'X\_'"

#### **Development Practices**

### Standards for OSS

Everything discussed in the open. Every convention and process documented.

# Development guide

http://scikit-learn.org/dev/developers/contributing.

#### **Contains:**

- API details
- Bug report guidelines
- PR guidelines
- Reviewing guidelines
- How to find issues
- Details of CI

# Deprecations / backward compatibility

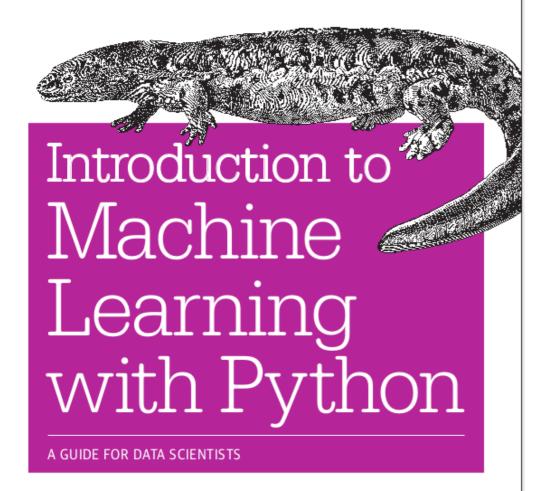
Don't change any behavior (except bug fixes)

(my personal) Roadmap

#### Make simple things simple again!

Missing Values
Categorical Variables
Plotting Tools
Pandas Integration
Feature Names

#### O'REILLY'



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